

GenCore version 5.1.3  
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OM nucleic - nucleic search, using sw model

Run on: March 30, 2003, 02:49:33 ; Search time 25.5808 Seconds

8283.243 million cell updates/sec

Title: US-09-988-971-1\_COPY\_694\_942

Sequence: 1 tggctgtatgagggcctgag.....aggccctggtgaccattac 249

Scoring table: IDENTITY\_NUC

Searched: 574371 seqs, 425486471 residues

Total number of hits satisfying chosen parameters: 1148742

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Minimum DB seq length: 0
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100

Post-processing: Minimum Match 0%

### Listing first 45 summaries

Database : Published Applications NA:\*

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1: /cgn2_6/p/odata/1/pubnba/us07 PUBCOMB.seq.*
2: /cgn2_6/p/odata/1/pubnba/pct_ NEW PUB.seq.*
3: /cgn2_6/p/odata/1/pubnba/us06 NEW PUB.seq.*
4: /cgn2_6/p/odata/1/pubnba/us06 PUBCOMB.seq.*
5: /cgn2_6/p/odata/1/pubnba/us07 NEW PUB.seq.*
6: /cgn2_6/p/odata/1/pubnba/pctus PUBCOMB.seq.*
7: /cgn2_6/p/odata/1/pubnba/us08 NEW PUB.seq.*
8: /cgn2_6/p/odata/1/pubnba/us08 PUBCOMB.seq.*
9: /cgn2_6/p/odata/1/pubnba/us09 NEW PUB.seq.*
10: /cgn2_6/p/odata/1/pubnba/us09 PUBCOMB.seq.*
11: /cgn2_6/p/odata/1/pubnba/us10 NEW PUB.seq.*
12: /cgn2_6/p/odata/1/pubnba/us10 PUBCOMB.seq.*
13: /cgn2_6/p/odata/1/pubnba/us50 NEW PUB.seq.*
14: /cgn2_6/p/odata/1/pubnba/us50 PUBCOMB.seq.*

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Pred. NO. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	199	79.9	763	10	US-09-967-550-953	Sequence 953, App
2	100.2	40.2	2015	10	US-09-954-456-1993	Sequence 1893, App
3	96	38.6	444	10	US-09-867-550-951	Sequence 951, App
4	89	35.7	1911	10	US-09-917-800-1611	Sequence 1611, App
5	87.6	35.2	2354	10	US-09-867-7688-300	Sequence 300, App
6	83	33.5	2665	10	US-09-967-550-1915	Sequence 1915, App
7	80.4	32.3	2875	10	US-09-954-456-499	Sequence 499, App
8	80.4	32.3	3756	12	US-10-002-600-931	Sequence 91, App
9	78.4	31.5	2451	10	US-09-771-1611-4	Sequence 4, App
10	73	29.3	1609	10	US-09-771-1611-30	Sequence 30, App
11	73	29.3	1995	9	US-09-771-1611-31	Sequence 31, App
12	58.2	14.2	486	9	US-09-796-692-7419	Sequence 7419, App
13	47	14.9	761	10	US-09-910-0-943-616	Sequence 616, App
14	35.4	18.2	920	9	US-09-864-480-127	Sequence 127, App
15	34	13.7	2770	9	US-09-977-261-5	Sequence 5, App
16	34	13.7	2770	9	US-09-977-261-5	Sequence 5, App
17	34	13.7	2770	9	US-09-977-261-5	Sequence 5, App
18	34	13.7	2863	10	US-09-954-456-1611	Sequence 1611, App
19	34	13.7	7607	10	US-09-982-610-19	Sequence 19, App

C	20	34	13.7	13873	9	US-09-764-668-1282	App	Sequence 1282, App
C	21	32	12.9	285	10	US-09-967-701-5885	App	Sequence 5885, App
C	22	31	12.4	366	10	US-09-950-063-40	App	Sequence 40, App
C	23	31	12.4	18400	10	US-09-901-151-3	App	Sequence 3, App
C	24	31	12.4	30365	10	US-09-825-414-1	App	Sequence 1, App
C	25	30.8	12.4	1552	10	US-09-925-297-226	App	Sequence 226, App
C	26	30.6	12.3	2888	9	US-09-822-946-209	App	Sequence 209, App
C	27	30.4	12.2	17849	9	US-10-092-154-144	App	Sequence 115, App
C	28	30.4	12.2	17849	9	US-09-764-847-1315	App	Sequence 1315, App
C	29	30.4	12.2	17862	9	US-10-092-154-1313	App	Sequence 1313, App
C	30	30.4	12.1	17862	10	US-09-764-847-1313	App	Sequence 1313, App
C	31	30.2	12.1	1725	9	US-10-103-196-7	App	Sequence 7, App
C	32	30.2	12.1	1749	9	US-10-103-196-13	App	Sequence 13, App
C	33	30	12.0	1646	9	US-10-016-634-94	App	Sequence 94, App
C	34	29.8	12.0	1058	10	US-09-880-107-2180	App	Sequence 2180, App
C	35	29.8	12.0	1425	9	US-10-102-806-149	App	Sequence 149, App
C	36	29.8	12.0	2382	10	US-09-964-824A-572	App	Sequence 572, App
C	37	29.6	11.9	596	9	US-10-023-380-931	App	Sequence 931, App
C	38	29.6	11.9	596	10	US-09-922-217-931	App	Sequence 931, App
C	39	29.6	11.9	596	10	US-09-833-263-931	App	Sequence 931, App
C	40	29.6	11.9	1668	10	US-09-925-301-208	App	Sequence 208, App
C	41	29.6	11.9	1834	10	US-09-948-094-1	App	Sequence 1, App
C	42	29.6	11.9	1834	10	US-09-880-107-2214	App	Sequence 2214, App
C	43	29.6	11.9	1834	10	US-09-967-768A-141	App	Sequence 141, App
C	44	29.4	11.8	900	9	US-10-014-64A-282	App	Sequence 282, App
C	45	29.4	11.8	1967	10	US-09-838-523-3	App	Sequence 3, App

## ALIGNMENTS

## RESULT 1

Sequence 953, Application US/09867550

GENERAL INFORMATION

APPLICANT: Mehraban, Fuad,

APPLICANT: Law, Debbie

TITLE OF INVENTION: NO. US200200822C

FILE REFERENCE: 21402-013 (Cura-313)

CURRENT FILING DATE: 2001-09-20

PRIOR FILING DATE: 2000-05-30

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; SOFTWARE: FastSEQ for windows Version 1.0

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LENGTH: 763

ORGANISM: Homo sapiens

1. **Introduction**

Best Local Similarity 100.0%; Prec

[illegible]

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**CONCLUSIONS**

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/ PRIOR FILING DATE: 2000-09-27
/ NUMBER OF SEQ ID NOS: 2276
/ SOFTWARE: ParentIn version 3.0.0
/ SEQ ID NO 499
/ LENGTH: 2665
/ TYPE: DNA
/ ORGANISM: Homo sapiens
US-09-954-456-499

```

Query Match	32.3%	Score 80.4	DB 10	Length 2665
Best Local Similarity	62.1%	Pred. No. 1.3e-16		
Matches 154; Conservative	0	Mismatches 76	Indels 18	Gaps 1

Qy 1 TGGCTGTAAGAGGAGCTTACGACGAGGAGAAAGAGGAACTGCTGTATTACCTGGAGAC 60  
Db 291 TGGCTGTTAGAGGAGCTTGGACAGAGACAAGAGCCGAGAGCTGTGCACCTGCCAGACACA 350  
Qy 61 CCGAGAGGGGCTTTCTCTATCCGAGAGACGACAGACAGAGAGGCTCTTACTCTCTGACA 120  
Db 351 AAGGTGGGCTCTTATGATGATGAGAGAGATGAGAGACAGAGAAAGGTTTACTACTGTGCG 410  
Qy 121 GTCCGCTCTAGGCCGCCCTGATCCTGGAGACGGATATGAGACATACAGATATCACTGCTTT 180  
Db 411 GTGAGACACAG-----GCAAGTAAGACATTACCGGATTTTTCGCTCG 452  
Qy 181 GACATAGGCTGTGCTGATCATCTTACACGCCCTCACTTTCCTCTACTCCAGCCCTGTG 240  
Db 453 CGGAACAACGTGATCACTATTTCCCGAGGCTCACTTCCAGTGTGAGAGACCTGTGG 512  
Qy 241 GACCATTA 248  
Db 513 AACCACTA 520

```

RESULT 8
US-10-002-600-91
Sequence 91, Application US/10002600
Patent No US20020137077A1
GENERAL INFORMATION:
APPLICANT: Hopkins, Christopher M.
APPLICANT: Peterson, David P.
APPLICANT: Cocks, Benjamin G.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: GENES REGULATED IN AC
FILE REFERENCE: PA-0042 US
CURRENT APPLICATION NUMBER: US/10/002,600
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: 60/243,521
PRIOR FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 116
SOFTWARE: PERL Program
SEQ ID NO 91
LENGTH: 3756
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Template ID: 059263.1
US-10-002-600-91

```

Query Match	32.3%	Score 80.4;	DB 12;	Length 3756;
Best Local Similarity	62.1%;	Pred. No. 1.3e-16;		
Matches 154; Conservative	0;	Mismatches 76;	Indels 18;	Gaps 1

QY	1	TGGCTGTATATAGAGCCCTAGACAGAGAGAAACAGAGAACTGTGCTTTTAACTGGGAAC	60
Db	1365	TGGCTGTTTAAAGGACCTGGGACAGACAAAGCCGAGGCTGTGAGTGTCCAGACACA	1424
QY	61	CCTGAGAGGAGCCTTCCCTATACCGGAGACCCAGACAGAGAGAGCTTATCTCTGTCA	120
Db	1425	AAGGTGGCTCTTTCATATATACAGAGAGATGAGACACAGAAAGGTTTTTACTCACTGTGC	1488
QY	121	GTGGCGCTCAGCGCCCTGCATCTGTGGACCGGATACAGACTTACAGATCCACTGCTTT	180

Db	1485	GTGAGACAAAG	-----	GCAGGTAAAGCATTAACCGCATTTTCCGCTCG	1526
OY	181	GACATGCGCTGCGCTGATCATCTCACGCGCCTCACTTCCCTCACTCCAGGCGCCCTG	240		
Db	1527	CCCAACACCTGGATCACTATTTCCCGAGGCTCACTTCCAGTGCCTGGAGGACCTGGTG	1586		
OY	241	GACCTTAA	248		
Db	1587	AACGACCTA	1594		

```

RESULT 9
US-09-771-161A-4
; Sequence 4, Application US/0977161A
; Patent No. US20020110811A1
; GENERAL INFORMATION:
;   APPLICANT: LEVINE, et al.
;   TITLE OF INVENTION: VARIANTS OF PROTEIN KINASES
;   FILE REFERENCE: 802620-2005.1
;   CURRENT APPLICATION NUMBER: US/09/771,161A
;   CURRENT FILING DATE: 2001-01-26
;   PRIOR APPLICATION NUMBER: 09/724,676
;   PRIOR FILING DATE: 2000-11-28
;   PRIOR APPLICATION NUMBER: 136776
;   PRIOR FILING DATE: 2000-06-15
;   PRIOR APPLICATION NUMBER: 135519
;   PRIOR FILING DATE: 2000-04-12
;   NUMBER OF SEQ ID NOS: 273
;   SOFTWARE: PatentIn version 3.0
;   SEQ ID NO: 4
;   LENGTH: 2451
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;     NAME/KEY: -
;     LOCATION: (1)..(2451)
;   OTHER INFORMATION: "n" can be any nucleotide 'a', 'c', 'g' or 't'
US-09-771-161A-4

```

Query Match	31.5%	Score 78.4	DB 10	Length 2451
Best Local Similarity	60.1%	Pred. No. 5	6e-16	
Matches 149	Conservative 0	Mismatches 96	Indels 3	Gaps 1
QY	1	TGCGCTGATGAGAGCGCTGTGACAGGGAGAAACCAAGAGAACTGTGTTGTTACCTGGAGAC	60	
DB	958	TGTTTCTTTAGATCAACAGGGGTGCGAAGAGAGCGTGAAGGCACTTTCTGTGCTCAATCAAC	1017	
QY	61	CCTGAGAGGGACCTTCTCTCATCCGGGAGAGCAGACAGGAGAGGCTTTACTCTGTGCA	120	
DB	1018	AAGCGCGGCTCTTTCTTTATCAAGAGAGTGTAAACCAAGAGTGCTTCTTCCCTGTCT	1077	
QY	121	GTCGCGCTCAGCGCGCTGTGATCTGTGGACCGGATCAGACACTACAGAGATCACTGCTCTT	180	
DB	1078	GTGAAGS--ATGTACCACCCAGGGGAGGCTGATCAACACTATTAAGATCGCGTGGCTG	1134	
QY	181	GACATATGCTGGCTGTATCATCTACCGGGCTCACTTCCCTCACTCAGAGCCTGTG	240	
DB	1135	GATGAAGGGGGCTACTACATCTCCCGGATCATCTTCCCTGTGCTCCAGGCGCTGTG	1194	
QY	241	GACCATTA	248	
DB	1195	CAGCACTA	1202	

RESULT 10  
 US-09-771-161A-30  
 ; Sequence 30, Application US/09771161A  
 ; Patent No. US20020110811A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: LEVINE, et al.  
 ; TITLE OF INVENTION: VARIANTS OF PROTEIN KINASES  
 ; FILE REFERENCE: 802620-2005.1

CURRENT APPLICATION NUMBER: US/09/771,161A  
CURRENT FILING DATE: 2001-01-26  
PRIOR APPLICATION NUMBER: 09/724,676  
PRIOR FILING DATE: 2000-11-28  
PRIOR APPLICATION NUMBER: 136776  
PRIOR FILING DATE: 2000-06-15  
PRIOR APPLICATION NUMBER: 135619  
PRIOR FILING DATE: 2000-04-12  
NUMBER OF SEQ ID NOS: 273  
SOFTWARE: PatentIn version 3.0  
SEQ ID NO 30  
LENGTH: 1609  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-09-771-161A-30

Query Match 29.3%; Score 73; DB 10; Length 1609;  
Best Local Similarity 55.8%; Pred. No. 2,9e-14;  
Matches 139; Conservative 0; Mismatches 110; Indels 0; Gaps 0;

Oy 1 TGGCTGTATGAGGCGCTGAGGAGGAGAAAGCAGAGGAACTGCTGTGTTACTCTGGAAAC 60  
Db 1023 TGGTACTTTGGAAGAACTGGCCGAAAGATGCTGAGGAGAGCTATGTGCTTTGGAAAC 1082  
Oy 61 CTTGAGAGGCGCTTCTTCATCCGAGAGCCAGACCCAGAGAGCTTTACTCTGTCTCA 120  
Db 1083 CCAAGAGGAGCTTTCTTATCCGAGAGTAAACCCAGAGGAGCTTATTCCTTTCT 1142  
Oy 121 GTCCGCTCAGCGCCCTGCATCTTGAGCGGATGACACACTACAGATCCACTGCTT 180  
Db 1143 ATCCGTGATGGATGATGATGAAAGAGACCATGTCAACATATTAATTCGAAACTT 1202  
Oy 181 GACATGCTGCTGTACATCTACCCGCGCTCACCTCCCTCCTCCAGCCCTGGG 240  
Db 1203 GACAAATGGTGAATCACTACATTACACCCGCGCCAGTTTGAACACTTACAGACTGTA 1262  
Oy 241 GACCATTTAC 249  
Db 1263 CACCATTTAC 1271

RESULT 11  
US-09-771-161A-31  
Sequence 31, Application US/09771161A  
Patent No. US20020110811A1  
GENERAL INFORMATION:  
APPLICANT: LEVINE, et al.  
TITLE OF INVENTION: VARIANTS OF PROTEIN KINASES  
FILE REFERENCE: 802620-2005.1  
CURRENT APPLICATION NUMBER: US/09/771,161A  
PRIOR FILING DATE: 2001-01-26  
PRIOR APPLICATION NUMBER: 09/724,676  
PRIOR FILING DATE: 2000-11-28  
PRIOR APPLICATION NUMBER: 136776  
PRIOR FILING DATE: 2000-06-15  
PRIOR APPLICATION NUMBER: 135619  
PRIOR FILING DATE: 2000-04-12  
NUMBER OF SEQ ID NOS: 273  
SOFTWARE: PatentIn version 3.0  
SEQ ID NO 31  
LENGTH: 1995  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: -  
LOCATION: (1)..(1995)  
OTHER INFORMATION: "n" can be any nucleotide 'a', 'c', 'g' or 't'  
US-09-771-161A-31

Query Match 29.3%; Score 73; DB 10; Length 1995;  
Best Local Similarity 55.8%; Pred. No. 3e-14;  
Matches 139; Conservative 0; Mismatches 110; Indels 0; Gaps 0;

Oy 1 TGGCTGTATGAGGCGCTGAGGAGGAGAAAGCAGAGGAACTGCTGTGTTACTCTGGAAAC 60  
Db 1140 TGGTACTTTGGAAGAACTGGCCGAAAGATGCTGAGGAGAGCTATGTGCTTTGGAAAC 1199  
Oy 61 CTTGAGAGGCGCTTCTTCATCCGAGAGCCAGACCCAGAGAGCTTTACTCTGTCTCA 120  
Db 1200 CCAAGAGGAGCTTTCTTATCCGAGAGTAAACCCAGAGGAGCTTATTCCTTTCT 1259  
Oy 121 GTCCGCTCAGCGCCCTGCATCTTGAGCGGATGACACACTACAGATCCACTGCTT 180  
Db 1260 ATCCGTGATGGATGATGATGAAAGAGACCATGTCAACATTAATTCGAAACTT 1319  
Oy 181 GACATGCTGCTGTACATCTCACCCGCGCTTACCTCCCTACCTCCAGGCGCTGGG 240  
Db 1320 GACAAATGGTGAATCACTACATTACACCCGCGCCAGTTTGAACACTTACAGACTGTA 1379  
Oy 241 GACCATTTAC 249  
Db 1380 CACCATTTAC 1388

RESULT 12  
US-09-796-692-7419  
Sequence 7419, Application US/09796692  
Publication No. US20020198362A1  
GENERAL INFORMATION:  
APPLICANT: Gaiger, Alexander  
APPLICANT: Algate, Paul A.  
APPLICANT: Mannion, Jane  
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DETECTION, DIAGNOSIS AND THERAPY  
FILE REFERENCE: 2077,001200  
CURRENT APPLICATION NUMBER: US/09/796,692  
CURRENT FILING DATE: 2001-03-01  
PRIOR APPLICATION NUMBER: 60/186,126  
PRIOR FILING DATE: 2000-03-01  
PRIOR APPLICATION NUMBER: 60/190,479  
PRIOR FILING DATE: 2000-03-17  
PRIOR APPLICATION NUMBER: 60/200,545  
PRIOR FILING DATE: 2000-04-27  
PRIOR APPLICATION NUMBER: 60/200,303  
PRIOR FILING DATE: 2000-04-28  
PRIOR APPLICATION NUMBER: 60/200,779  
PRIOR FILING DATE: 2000-04-28  
PRIOR APPLICATION NUMBER: 60/200,999  
PRIOR FILING DATE: 2000-05-01  
PRIOR APPLICATION NUMBER: 60/202,084  
PRIOR FILING DATE: 2000-05-04  
PRIOR APPLICATION NUMBER: 60/206,201  
PRIOR FILING DATE: 2000-05-22  
PRIOR APPLICATION NUMBER: 60/218,950  
PRIOR FILING DATE: 2000-07-14  
PRIOR APPLICATION NUMBER: 60/222,903  
PRIOR FILING DATE: 2000-08-03  
PRIOR APPLICATION NUMBER: 60/223,416  
PRIOR FILING DATE: 2000-08-04  
PRIOR APPLICATION NUMBER: 60/223,378  
PRIOR FILING DATE: 2000-08-07  
NUMBER OF SEQ ID NOS: 9597  
SOFTWARE: FastSeq for Windows Version 3.0  
SEQ ID NO 7419  
LENGTH: 486  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-09-796-692-7419

Query Match 23.4%; Score 58.2; DB 9; Length 486;  
Best Local Similarity 57.4%; Pred. No. 1.5e-09;  
Matches 105; Conservative 0; Mismatches 78; Indels 0; Gaps 0;  
Oy 67 GGGGCTTCTCTATCCGAGAGCCAGACCCAGAGAGCTTTACTCTCTGCTGCTCCG 126  
Db 7 GGAGCTTCTCTATTAGAGAACTGAACATTAAAGAGAGCTTCTCTGCTGCTGCA 66

QY	127	CTGAGCGCGCCGAGTCTCTGGAGACGGATGACATTAAGATTCACCTCCCTTGACAT	186
DB	67	GATTTGAGCCCTGGAGCATGCTATGTATTATAGACACTCAAAATTTAAGATCTGGATTAAT	128
QY	187	GAGCTGCTGTACATCTACACGGCGCTCACTTCCCTCACTCCAGCGCCCTGGTGACCAT	246
DB	127	GGGGGCTATTACATCTCTCCAGAAATCACTTTCCCTGATGACGACATGATTAAACAT	186
QY	247	TAC	249
DB	187	TAC	189

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RESULT 13
US-09-910-943-616
: Sequence 616, Application US/09910943
: Patent No. US20020081610A1
: GENERAL INFORMATION:
: APPLICANT: Hemmati-Brivianlou, Ali
: APPLICANT: Altman, Curtis
: TITLE OF INVENTION: Assays and Materials for Embryonic Gene Expression
: FILE REFERENCE: 7529/1G148US1
: CURRENT APPLICATION NUMBER: US/09/910,943
: CURRENT FILING DATE: 2001-07-23
: NUMBER OF SEQ ID NOS: 742
: SOFTWARE: PatentIn version 3.1
: SEQ ID NO. 616
: LENGTH: 761
: TYPE: DNA
: ORGANISM: Xenopus laevis
: FEATURE:
: NAME/KEY: misc_feature
: LOCATION: (1)..(761)
: OTHER INFORMATION: n may be a or g or c or t/u
US-09-910-943-616

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RESULT 14  
US-09-986-480-127/c  
Sequence 127, Application US/09986480  
Publication No. US20030027999A1  
GENERAL INFORMATION:  
APPLICANT: Rosen et al.  
TITLE OF INVENTION: 143 Human Secreted Proteins  
FILE REFERENCE: PSS00p1  
CURRENT APPLICATION NUMBER: US/09/986,480  
CURRENT FILING DATE: 2001-11-08  
PRIOR APPLICATION NUMBER: PCT/US00/12786  
PRIOR FILING DATE: 2000-05-11  
PRIOR APPLICATION NUMBER: US 60/134,068  
PRIOR FILING DATE: 1999-05-13  
NUMBER OF SEQ ID NOS: 46

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; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 127
; LENGTH: 920
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-986-480-127

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Query Match	14.2%	Score 35.4;	DB 9;	Length 920;
Best Local Similarity	48.3%;	Pred. NO. 0.038;		
Matches 99; Conservative	0;	Mismatches 106;	Indels 0;	Gaps 0

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RESULT 15
US-09-977-260-5
Sequence 5, Application US/09977260
Publication No. US20020192790A1
GENERAL INFORMATION:
APPLICANT: ULLRICH, AXEL
APPLICANT: GISHIZKY, MICHAEL
APPLICANT: SURES, IRMINARD
TITLE OF INVENTION: NOVEL MEGAKARYOCYTIC PROTEIN TYROSINE KINASES
FILE REFERENCE: 038602/1,660
CURRENT APPLICATION NUMBER: US/09/977,260
CURRENT FILING DATE: 2001-10-16
PRIOR APPLICATION NUMBER: 08/232,545
PRIOR FILING DATE: 1994-04-22
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
LENGTH: 2770
TYPE: DNA
ORGANISM: Unknown Organism
FEATURE:
NAME/KEY: CDS
LOCATION: (366)..(1880)
FEATURE:
OTHER INFORMATION: Description of Unknown Organism: Megakaryocyte
OTHER INFORMATION: Kinase 3
US-09-977-260-5

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Search Completed: March 30, 2003, 16:31:13  
Job time : 31.5808 secs

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